

INHIBITORS OF DIPEPTIDYL-AMINOPEPTIDASE TYPE IV Abstract of the Disclosure

An inhibitory compound having the structure: Group I - Group II. Group I has the structure:

$$H = \begin{bmatrix} H & O & & & O \\ & & & & & \\ & & & & \\ NH' - C - C - N - C - C \end{bmatrix} \begin{bmatrix} H \\ & & \\ NH' - C \\ & & \\ R \end{bmatrix}$$

$$R1 - C - Y$$

$$R2$$

where H represents a hydrogen; C represents a carbon; O represents an oxygen; N represents a nitogen; each R, independently, is chosen from the group consisting of the R groups of an amino acid, including proline; each broken line, independently, represents a bond to an H or a bond to one R group, and each H' represents that bond or a hydrogen; and p is an integer between 0 and 4 inclusive.

Alternatively Group I has the structure:

$$G1 \left\{ \begin{array}{c} G2 \\ | \\ C \\ | \\ G3 \end{array} \right\}_{n}$$

where n is between 0 and 3 inclusive, each G2 and G3 independently is H or C1 - 3 (one to three carbon atoms) alkyl, G1 is NH3 (H3 represents three hydrogens),

(H2 represents two hydrogens), or

NG4, where G4 is
$$C - G5$$

where G5 and G6 can be NH, H, or C1 - 3 alkyl or alkenyl with one or more carbons substituted with a nitrogen. G1 bears a charge, and G1 and Group II do not form a covalently bonded ring structure at pH 7.0. Group I may also have the structure:

where one or two of the a, b, c, d, e, and f group is N, and the rest are C, and each S1 - S6 independently is H or C1 - C3 alkyl. Group I may also include a five membered unsaturated ring having two nitrogen atoms. Group II has the structure:

where T is a group of the formula:

D2

- B - D1, where each D1 and D2, independently, is a hydroxyl group or a group which is capable of being hydrolysed to a hydroxyl group in aqueous solution at physiological pH; a group of the formula:

where G is either H, fluorine (F) or an alkyl group containing 1 to 20 carbon atoms and optional heteroatoms which can be N, S (sulfur), or O; or a phosphonate group of the formula:

where each J, independently, is O-alkyl, N-alkyl, or alkyl. Each O-alkyl, N-alkyl or alkyl includes 1 - 20 carbon atoms and, optionally, heteroatoms which can be N, S, or O. T is generally able to form a complex with the catalytic site of a DP IV.

and each R1, R2, R3, R4, R5, R6, R7, and R8, separately is a group which does not significantly interfere with site specific recognition of the inhibitory compound by DP IV, and allows a complex to be formed with DP IV.

In another aspect, the invention features an inhibitor of DP-IV, having the structure:

$$\begin{bmatrix} A & -N & -C & -C & -A' & -N & -C & -B \\ -CH_2 & -C$$

wherein m is an integer between 0 and 10, inclusive; A and A' are L-amino acid residues such that the A in each repeating bracketed unit can be a different amino acid residue; the C bonded to B is in the L-configuration; the bonds between A and N, A and C, and between A and N are peptide bonds; and each X¹ and X² is, independently, a hydroxyl group or a group capable of being hydrolysed to a hydroxyl group at physiological pH.

APPL2378.BOS